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Sequence Listing was accepted.

See attached Validation Report.

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Reviewer: Anne Corrigan

Timestamp: [year=2008; month=12; day=4; hr=7; min=24; sec=12; ms=815;]

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Application No: 09540245 Version No: 1.0

Input Set:

Output Set:

Started: 2008-12-02 10:18:59.302
Finished: 2008-12-02 10:19:02.950
Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 648 ms
Total Warnings: 14
Total Errors: 1
No. of SeqIDs Defined: 20
Actual SeqID Count: 20

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W 402	Undefined organism found in <213> in SEQ ID (2)
W 402	Undefined organism found in <213> in SEQ ID (3)
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W 402	Undefined organism found in <213> in SEQ ID (5)
E 341	'Xaa' position not defined SEQID (5) POS (153)
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W 402	Undefined organism found in <213> in SEQ ID (19)
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<110> APPLICANT: Goodman, Corey
Kid, Thomas
Brose, Katja
Tessier-Lavigne, Marc
<120> TITLE OF INVENTION: Modulating Robo: Ligand Interactions
<130> FILE REFERENCE: B98-031-3

<140> CURRENT APPLICATION NUMBER:09540245
<141> CURRENT FILING DATE:2000-03-31
<150> PRIOR APPLICATION NUMBER: 60/065,544
<151> PRIOR FILING DATE: 1997-11-14
<150> PRIOR APPLICATION NUMBER: 60/081,057
<151> PRIOR FILING DATE: 1998-04-07
<160> NUMBER OF SEQ ID NOS: 20
<170> SOFTWARE: PatentIn Ver. 2.0

<210> SEQ ID NO 1
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<212> TYPE: DNA
<213> ORGANISM: human
<220> FEATURE:
<221> NAME/KEY: CDS
<222> LOCATION: (1)..(4755)
<400> SEQUENCE: 1

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Ser Cys Ser Gly Ser Thr Val Asp Cys His Gly Leu Ala Leu Arg Ser	
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gtg ccc agg aat atc ccc cgc aac acc gag aga ctg gat tta aat gga	192
Val Pro Arg Asn Ile Pro Arg Asn Thr Glu Arg Leu Asp Leu Asn Gly	
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cta aga gtt ctt cag ctt atg gag aat aag att agc acc att gaa aga	288
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Gly Ala Phe Gln Asp Leu Lys Glu Leu Glu Arg Leu Arg Leu Asn Arg	
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Asn His Leu Gln Leu Phe Pro Glu Leu Leu Phe Leu Gly Thr Ala Lys	
115 120 125	
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Leu Tyr Arg Leu Asp Leu Ser Glu Asn Gln Ile Gln Ala Ile Pro Arg	
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Ala Ser Phe Asn His Met Pro Lys Leu Arg Thr Phe Arg Leu His Ser	
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Ala Asn Lys Ile Asn Cys Leu Arg Val Asp Ala Phe Gln Asp Leu His	
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Gln Asn Pro Phe Ile Cys Asp Cys His Leu Lys Trp Leu Ala Asp Tyr	
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Leu His Thr Asn Pro Ile Glu Thr Ser Gly Ala Arg Cys Thr Ser Pro	
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ccc	tcc	aaa	aaa	ttt	acc	tgt	caa	ggc	cct	gtg	gat	gtc	aat	att	cta	2736
Pro	Ser	Lys	Lys	Phe	Thr	Cys	Gln	Gly	Pro	Val	Asp	Val	Asn	Ile	Leu	
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Ala	Lys	Cys	Asn	Pro	Cys	Leu	Ser	Asn	Pro	Cys	Lys	Asn	Asp	Gly	Thr	
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